

UNIVERSITY OF CALIFORNIA

DEPARTMENT OF PHYSICS
BERKELEY 4, CALIFORNIAMay 27th

Dear Francis,

Thank for your letter. Now I have completely shifted to RNA. When Jim was in the east I visited Pasadena, and together with Alek Rich and Leslie Orgel (you must know him, he is from Oxford) we devised a new code in which an a.a. is determined by three consecutive bases, the central one being a "major determinant", with two neighbours "only collaborating". Something like that:

~~Does not work~~ Ad Gc Ur Ad Gu Ur ect J+ looked at first very promising,
Does not work Ad Gc Ur Ad Gu Ur ect Ala Ser Glo ect but after several days of hard work (using about 70 known exact analogies three peptides) Dick Finemann ~~succes~~ succeeded to show that no solution exist. I am now trying a new code recently suggested by Edward Teller in which each new a.a. is determined by two bases and the preceding aa. (This has a direction!). Do not

~~not wait.~~ Not wait. Base Direction of growth ↑ <span style="color: red; position: absolute;